

The motion consists of shimmering points which are perpendicular to the lines of the distal stimulus. Thus, with a radiating pattern they are circular, and can be seen more readily within the orange annuli. If a white surface is observed following fixation on the pattern for around 30 seconds then scintillating dots will seem to be streaming in a direction orthogonal to the previously presented lines.

Regular geometrical patterns produce a range of visual effects; some are based on the optics of the eye (like the spokes) and others (such as the scintillations) could be caused by visual processing in the brain. For example, Purkinje's radiating lines illusion has been examined with the aid of neural imaging. Positron emission tomography (PET) scans indicate that a pattern of radiating lines causes the blood flow in V5 — the area of the brain that is involved in motion processing — to increase. Alternative interpretations, however, relate to eye movement instabilities during fixation. It is likely that several phenomena are induced by these patterns, with a different explanation for each one.

When we know that an image is distorted before it reaches the retina — that the proximal stimulus differs from the distal stimulus — we consider that we have some interpretation of the illusion. Thus, the bent-stick distortion is due to differential refraction through air and water. But in general, explanations for the illusion are sought in higher levels of visual processing. Such illusions might provide an index of neural function, which I have called a 'neuro-sign'.

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This is the first in a short series of articles by Nicholas Wade, describing some common visual illusions.

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A bacterial antibiotic resistance gene with eukaryotic origins

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The rapid emergence of antibiotic-resistant pathogens is mainly due to the horizontal transfer of resistance genes among different bacterial strains or species. No transfers of antibiotic resistance genes have been found to occur between bacteria and eukaryotes, however [1]. Here, we show that a gene known to convey antibiotic resistance in bacteria was apparently transferred from an early evolved eukaryote.

Mupirocin, a topical agent widely used since 1985 against skin surface and nasal colonization by the Gram-positive bacterium *Staphylococcus aureus*, is the natural compound pseudomonic acid produced by the bacterium *Pseudomonas fluorescens* NCIB 10586 [2]. Pseudomonic acid inhibits protein synthesis by reversibly binding to the active site of isoleucyl-tRNA synthetase (IRS) [3,4]. Sensitivity of *S. aureus* to mupirocin use *in vitro* occurs at a minimal inhibitory concentration (MIC) range of 0.12 to 1.0 mg/L [5,6]. Moderate mupirocin resistance in *S. aureus*, broadly defined as an MIC range of 8 to 256 mg/L, appears to be mediated by different single amino-acid substitutions in IRS, either within or near an ATP-binding motif with the consensus amino-acid sequence KMSKS (in the single amino-acid code) [7]. First observed clinically in the late 1980s, this level of resistance has little impact on the topical use of mupirocin.

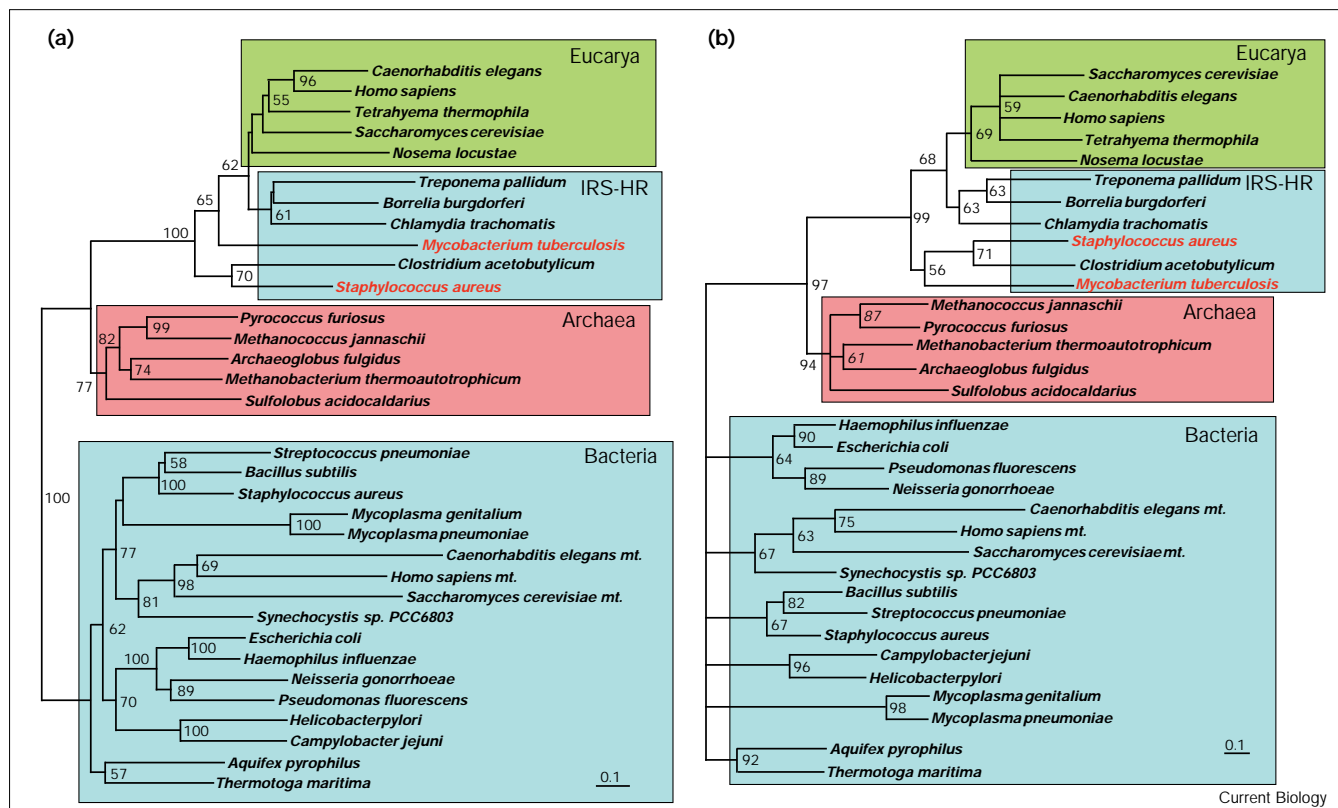
In the early 1990s, much higher levels of mupirocin resistance in *S. aureus* appeared in some clinical

settings, where estimated MICs exceed 512 mg/L [5]. Highly resistant strains have a second copy of the IRS gene (IRS-HR), located on an extrachromosomal plasmid [8], the product of which is highly divergent (34% amino-acid identity) from those of sensitive or moderately resistant IRS genes (IRS-S/MR) [9,10]. More recently, another Gram-positive bacterium, *Mycobacterium tuberculosis*, was shown to be highly resistant to mupirocin, and the product of its chromosome-located IRS gene had higher sequence similarity to *S. aureus* IRS-HR proteins than IRS-S/MR proteins [11].

Using the *S. aureus* IRS-HR sequence as a probe against public databases [12], we discovered four more bacterial IRS-HR-type genes. Phylogenetic analyses (Figure 1) show that all six bacterial IRS-HR proteins cluster significantly with eukaryotic IRS proteins that are also highly resistant to mupirocin [3,4]. The present IRS tree contradicts an earlier phylogeny, which showed the Bacteria, Archaea and Eucarya as separate monophyletic groups, with the latter two being sister groups [13]. We suggest that bacterial IRS-HR genes originated from eukaryotes through a horizontal gene transfer that occurred after the Eucarya and Archaea separated. A eukaryotic origin of bacterial IRS-HR genes is also supported by an insertion with the consensus sequence HYPFE, which is shared with eukaryotic IRS proteins but not bacterial IRS-S/MR proteins (Figure 2). Amino-acid compositions are significantly different between the IRS-HR and IRS-S/MR proteins of *S. aureus* (data not shown), further suggesting that the two genes originated from different organisms. A similar gene transfer has been suggested for glutamyl-tRNA synthetases, for which certain proteobacteria have a eukaryote-like gene [14,15].

The deep phylogenetic position of bacterial IRS-HR proteins relative to eukaryotic IRS proteins suggests that a recent transfer of the

Figure 1



Phylogenies of isoleucyl-tRNA synthetases (IRSs) using (a) neighbor-joining (NJ) [17] and (b) maximum likelihood (ML) [18] methods. Those bacterial IRSs suggested to be highly mupirocin resistant are indicated (HR), and those species in which it has been confirmed by experiments are in red. Trees are based on the non-gapped multiple sequence alignment of amino acids flanked by class 1 aminoacyl-tRNA synthetase signature motifs HIGH and KMSKS (438

amino acids). The NJ tree was generated with the programs PROTDIST (using the Dayhoff PAM120 option) and NEIGHBOR of the PHYLIP package [19]. Maximum log likelihood of the ML tree (using the program PUZZLE 3.0.1 [18] with the JTT option) was $-23,862.83 (\pm 514.99)$. Maximum parsimony (MP) analysis (using the program PAUP 3.1.1 [20]) also clustered IRS-HR sequences with eukaryotes with one minimal length tree (4295 steps) found after 100

random heuristic searches. Numbers represent the percent occurrence of branching points in 1000 bootstrap replications (NJ) or 1000 puzzling quartets (ML). Values less than 50% are not shown, and in the ML tree those nodes are collapsed. The scale bar represents an estimated 0.1 amino-acid substitutions per site. Mitochondrial targeted isoforms of eukaryotes are indicated by mt.

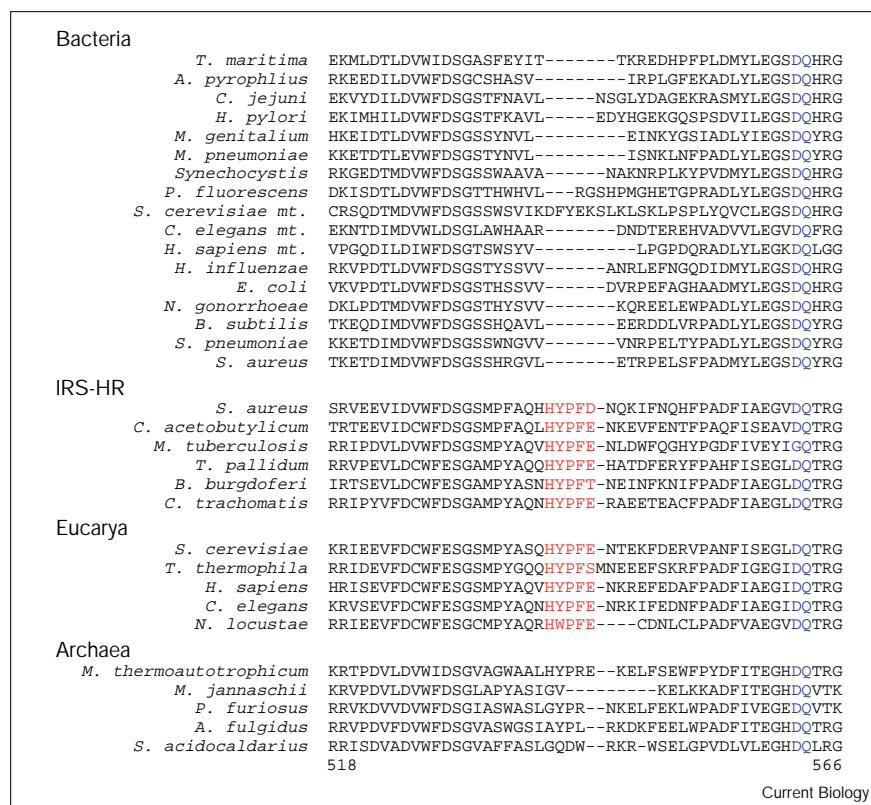
resistance gene from humans or animals as a consequence of intensive clinical use of mupirocin is an unlikely scenario. Plasmids mediating high-level resistance were detected in staphylococci isolated nearly twenty years before the clinical use of mupirocin [16]. The existence of mupirocin-sensitive *S. aureus* and plasmid-borne IRS-HR genes, however, indicates that IRS-HR is not fixed in the population. To reconcile these observations, we propose that a eukaryotic IRS gene was first transferred to an unknown bacterium shortly after the divergence of the

Eucarya and Archaea. The gene might have been transferred through several bacteria species and only recently was it passed to *S. aureus*.

Since mupirocin (as pseudomonic acid) is a natural product of some widely found strains of *P. fluorescens*, resistance to this compound might have conferred some competitive advantages to conspecific organisms, thus driving the fixation of IRS-HR genes in different bacterial species, though the possibility of random fixation cannot be excluded. To our knowledge, *Treponema pallidum*, *Borrelia burgdorferi*, *Clostridium acetobutylicum* and *Chlamydia*

trachomatis have not been tested for susceptibility to mupirocin, probably because these organisms are not involved in epidermal infections and several are intracellular, obligate human parasites. The present analysis, however, strongly suggests these organisms to be resistant to mupirocin, as their IRS proteins clearly cluster with IRS-HR proteins from species with proven resistance. Our study shows the possibility of horizontal transfers of antibiotic resistance genes from eukaryotes to prokaryotes and highlights the potential of phylogenetic analyses in medical studies.

Figure 2



Partial amino-acid sequence alignment showing the consensus amino-acid motif HYPFE (in red) uniquely shared among bacterial IRS-HR and eukaryotic IRS.

Numbering corresponds to the first amino acid in human IRS. The di-amino-acid motif DQ (blue) potentially functions in binding the activated amino acid isoleucine-AMP [21].

Supplementary material

An alignment of multiple IRS amino-acid sequences is published with this article on the internet.

Acknowledgements

We thank K. Coleman, R. Jarvest, E. Lawlor, A. Lupas, C. Perry and J. Poupard for their comments. J.Z. was supported by NIH and NSF research grants to M. Nei.

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A bacterial antibiotic resistance gene with eukaryotic origins [Correspondence]

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Current Biology 21 May 1998, 8:365–367

Alignment of multiple isoleucyl-tRNA synthetase (IRS) amino-acid sequences

Columns marked with * were used in the phylogenetic analyses. Abbreviations correspond to IRS sequences from the following species with their GenBank accession numbers or source: ITma (*Thermotoga maritima*, L37104), IApY (*Aquifex pyrophilus*, L37096), ICje (*Campylobacter jejuni*, U15295), IHpy (*Helicobacter pylori*, AE000642), ISau (*Staphylococcus aureus*, X74219), ISaR (*Staphylococcus aureus Mup R* gene, X75439), ITpa (*Treponema pallidum*, The Institute for Genomic Research, personal communication), IBbu (*Borrelia burgdorferi*, AE001181), ICTr (*Chlamydia trachomatis*, Chlamydia Genome Project, personal communication), IMge (*Mycoplasma genitalium*, U39718), IMPn (*Mycoplasma pneumoniae*, 2501010), IMtu (*Mycobacterium tuberculosis*, Z74020), IBsu (*Bacillus subtilis*, Z99112), ICac (*Clostridium acetobutylicum*, Genome Therapeutics personal communication), ISpn (*Streptococcus pneumoniae*,

The Institute for Genomic Research, personal communication), ISyn (*Synechocystis* sp. PCC6803, D90907), IPfl (*Pseudomonas fluorescens*, X80132), ISmt (*Saccharomyces cerevisiae* mitochondria (mt.), L38957), ICmt (*Caenorhabditis elegans* mt., Z81038), IHmt (*Homo sapiens* mt., assembled from NCBI dbEST), IHin (*Haemophilus influenzae*, U32777), IEco (*Escherichia coli*, D10483), INgo (*Neisseria gonorrhoeae*, University of Oklahoma's Advanced Center for Genome Technology, personal communication), ISce (*Saccharomyces cerevisiae*, M19992), ITth (*Tetrahymena thermophila*, M30942), IHsa (*Homo sapiens*, U04953), ICel (*Caenorhabditis elegans*, Z70310), INlo (*Nosema locustae*, L37097), IMth (*Methanobacterium thermoautotrophicum*, M59245), IMja (*Methanococcus jannaschii*, U67538), IPfu (*Pyrococcus furiosus*, L37105), IAfu (*Archaeoglobus fulgidus*, AE001061), ISac (*Sulfolobus acidocaldarius*, L37106).

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#ICje -----LIFASALLKSMV-----EKGLTSGEIQKELNAKEFEKLEAINPLNGRKS-----
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#ISau ---GEKYIIAEALSDAVAEAL--DWDKASIKLEKEYTGKELEYVVAQHPFLDRESL-----
#ISaR ---NEYYILATDLINS-----IITEKYEIIDTFSGSNLINLKYIPPFESDGLV-----
#ITpa ---DEHYILARSRLFY-----YPDSSAYRVVWEKRGELAGIRYRPLFSYPVFGQGPDPSPVQGD
#IBbu ---TKEEILILGSKKLN---SYDDENSITYIIEKFKGSKLEGIEYEPIFNYFLEQKDKGAF-----
#ICtr -ESGDEYILQGESLPRW-----FPDRESYEWIGQLSGKSLVGQSYEPLFPYFQDK-----
#IMge ---QKFVILEKLFVEVFTNKL---NWTNAIKLKKFKGSNLKNSSSYSHCFYNKVLP-----
#IMpn ---NEQYVVLASLFESLKTKF---GWTDAIQVQTISGSQQLQNTTYKHCLYDKVNP-----
#IMtu --DRRFVLAEARLAAYAREL-GEEPVVLTGYRGAELLGTRYLPPFAYFMDWPNF-----
#IBsu --EDRFVVASALVENVASAC-----GFDQYEVTRTVKGKDLENIIEAHPYGRDSL-----
#ICac --GEHLILAKALLTVLEGEY-----EVVSEFKGEKLLGMEYEQLFKFANPDKKAFY-----
#ISPn -GEARKFVVAEELTLSEKF-----GWADVQVLETYRGQELNHIVTEHPWDTAVEE-----
#ISyn SHIFHRSYLIVALDLVEKLS---ETFGVKLTVKVTLQGESLENTCYQHPLFDRVSP-----
#IPfl ---DRLLVLAEEEMVESCLA-----RYELQGSVIATATGSALELINFRHPFYDRLSP-----
#ISmt ---SELILVETGSIDKLGL---TTNSFETIKQFQGTHLNGLYYQNLVDDKVGRP-----
#ICmt -EINNPTSTFYVIASKLLEEFQKSSDRKCKVVGTVKPANLIGRRYKSCWHNELGLP-----
#IHmt ---GDLYVLAADKVASVAS-----TLETTFETISTLSGVDLENGTCSHPLIPDKAS-----
#IHin ---DERVILAAELVESVAK---AVGIEHIEILGSVKGDDLELSRFHHPFYDFTVP-----
#IEco ---QAVILAKDLVESVMQ---RIGVTDYTLGTVKGADVELLRFTHPFMGFDVP-----
#INGo ---KGKLVLAKDLAGALK---RYGFSDGIAILAETTGDKLENLHMNHPFLERDIP-----
#ISce -TRDRYFILLESLIKTLYK---KPKNEKYKIVEKIKGSDLVGLKYEPLFPYFAEQFH-----
#ITth -KTQEHYILAECRLPELYK---KDKDGYKILEKFKGSELVGREYEPLFPYFLSRKQ-----
#IHsa -ARGRLILMEARLSALYK-----LES DYELERFPGAYLKGKKYRPLFDYFLKCKE-----
#ICel -TTGIEYVLEERLGELKN-----DNLEVIEKLAGSQLKDLRYEPLFPYFAYMRE-----
#INlo -LKS AFFIMQRDRVNVYFK-----DAVVISEVKGHELLHLEHDQPFMYNHYRE-----
#IMth ---GETYIMAEALVEKVLG-----EEAEI IKTVRGSELEGLTYRHPLDEEVPCHR-----
#IMja DDKKEVWIIAEKLVEEVINKAKKFHNKNIKI IKKVKGKELEGIKYIHPLLEENERQKEFAE----
#IPfu NGREEYWILAKALVEKVLG-----DIGVKGEVVEEFKGKELEGLRYVHILMDEYPRQKEFRE----
#IAfu DGKVEYLILAKELADSVLG---KG DYDSWEVVETYLGEDLEGLEYEHPLADEVPLQK-----
#ISac -----DEILVIAKDRVKEL--MKEARIKEYKILRVYKGEELLGLEYEHPLADIVSAQS-----

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#ITma  -----RVILAD-YVSLETGTGCVHIAPGHGEEDYIYGHVQYGL-----PIVSPVDEEGRF-----
#IApy  ETLKNMWRIYPSEFVSLDTGTGLVHMAPGHGQEDYTVGK-RYNL-----EPYAPLDDSGRF-----
#ICje  -----LIMGE-HVLMDDGSGLVHTAPGHGEDDYIYACL-KYGI-----EVLMPVDDSGCYDET LRARLL
#IHpy  -----VALGE-HVGLDGTGAVHTAPGHGEEDYIYGL-RYNL-----EVLMSVDEKGCYDEGI IHNQLL
#ISau  -----VINGD-HVTTDAGTGCVHTAPGHGEDDYIVGQ-KYEL-----PVISPIDDKGVF-----
#ISaR  ---NAYYVVDGE-FVTNSEGTGIVHIAPAHGEDDYQLVL-ERDL-----DFLNVITREGVY-----
#ITpa  SEEGLFCTRVAD-FVSTEDGTGVVHVAPAFGEDDYEVFK-DAGI-----SIQCPLDAECRF-----
#IBbu  -----KVHTAD-YVTTDDGTGIVHIAPFGEEDYRILKK-HTNV-----DIIDPLDAECKF-----
#ICtr  KELEAFRILPAD-FIEESEGTGIVHMAPAFGEADFFACQ-EHNV-----PLVCPVDNQGCY-----
#IMge  -----VLMGI-HVVDNEGTGIVHSSPAFGIDDFYLCQ-KNKI-----KEVLISIDEKGVF-----
#IMpn  -----VLLGN-HVLCNEGTGLVHTSPAYGLDDFYLCQ-QNKL-----NEALVSLDEKGVF-----
#IMtu  -----QVLAD-FVTTDDGTGIVHMAPAYGEDDMVVAE-AVGI-----APVTPVDSKGRF-----
#IBsu  -----VMLGE-HVTTDAGTGCVHTAPGHGEDDFIIGQ-KYGL-----DVLCPVDEKGVN-----
#ICac  -----VVHGD-FVTLSDDGTGIVHIAPAYGEDDNMLGK-KYDL-----PLINLVNNEGKVF-----
#ISpn  -----LVILGD-HVTTDSGTGIVHTAPGFGEEDYNVGI-ANNL-----EVAVTVDER GIM-----
#ISyn  -----IVIGGD-YVTTESGTGLVHTAPGHGQEDYVVGQ-RYGL-----PILSPVDAAGNL-----
#IPfl  -----VYLAD-YVELGSGTGIVHCSPAYGVDDFVICK-KYGM-----VNDDIINPVQSN-----
#ISmt  -----LLHGA-HVTSGTGTGLVHTAPGHGQDDYILGI-QNGL-----EIYSPVDHQGRYQLN-----
#ICmt  -----IYEGP-HVTDTVGTGLVHTAFAGHFQDYDAI-SKGD-----RVESFVDSRGCYTRH-----
#IHmt  -----PLLPPNHVTMAKGTGLVHTAPAHGMEDYGVAS-QHNL-----PMDCLVDEDDGVF-----
#IHin  -----VILGD-HVTTDGGTGLVHTAPDHGLDDFYVGI-QYDL-----PMAGLVSNDDGKF-----
#IEco  -----AILGD-HVTLDAGTGAVHTAPGHGPDDYVIGQ-KYGL-----ETANPVGPDGTY-----
#INGo  -----MLNGE-HVTTDAGTGLVHTAPAHGLEDYAVCN-KYGI-----ELYNPVNAEGKY-----
#ISce  --ETAFRVISDD-YVTSDDSGTGIVHNAFAGEEDNAACL-KNGVISEDVLPNAIDDLGRF-----
#ITth  --DGCFRILAGD-FVTADAGTGIVHCAPFGDDDYKVS-ANNI IKPDDPPV-PVDENGHF-----
#IHsa  --NGAFTVLVDN-YVKEEGTGVVHQAPYFGAEDYRVCM-DFNI IRKDSLPCVPVDASGCF-----
#ICel  --ERNAFRVLNDTFVTSDDSGTGIVHNAFAGEEDYVGI-QYDL-----ANGVIAKDQKMICPVDESGKY-----
#INlo  --KGFFRVYHAD-FVSEIDGTGVVHCSPFGGEDDYKAMT-MPGLIKENELLPSPLDENGFR-----
#IMth  --DMEHRVILGD-HVTLTGEGTGCVHTAPGHGPEDFEIGK-EYGL-----PVFCPVDEAGVF-----
#IMja  -LENAHTVILGE-HVTLEGGTGLVHTAPGHGEEDFEVGI-KYNL-----PIYSPIDDEGKY-----
#IPfu  KYEWWHRVILAD-FVTLEEGTGLVHTAPGHGEEDFEVGI-KYGL-----PVYSPVDDQGKY-----
#IAfu  ---NWKHQVFFA-DFVTAENTGCVHIAPGHGVEDYELGV-EKGL-----EVFNPVDDRGRVY-----
#ISac  KINNHHKVLDDGGEAVTLQEGTGLVHSAPGHGDVDFEIGK-KYDM-----PVVMLVNDKGEF-----

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#ITma --TEEAGKY-KGMFIE-DANEVIEDLKRK-----GILVHASSITHSYPHCWRCCKPVIIFRATQWFI
#IApy --VEPAPEFIRGVRVF-DANKLI IALLKEK-----GYLVHEARIRHSYPHCWRCCKNPVIFRATPQWFI
#ICje -PSHLLLEEFI-GLHIF-KANEKILELLGEK-----LLHSSKFIHSYPFCWRTHKPVIYRATKQWFIL
#IHpy -----DESYL-GEHVF-KAQKRIIEQLGDS-----LLLEQEIHSYPHCWRTHKPVIYRATTQWFI
#ISau --TEEGGQF-EGMFYD-KANKAVTDLLTEK-----GALLKLDFITHSYPHDWRTKKPVIIFRATPQWFAS
#ISaR --NDRFPEL-VGNKAK-NSDIEI I KLLSKK-----QLLYKKQKYEHNYPHCWRCGNPLIYYAMEGWFIK
#ITpa --TAEVADY-QGLFVK-AADKAI IARVQKQ-----GALFRREQISHAYPHCWRCASPLIYRAVHSWFVA
#IBbu --TNQVKDF-KGLFVK-DADKKI IENLKL R-----NFLFKRENYLHRYPF CYRTNCPIIYRPISSWFVN
#ICtr --TAEVKDF-VGEYIK-SADKGIARRLKNE-----NKL FYQGTVRHRYPF CWRTDSPLIYKAVNSWFVA
#IMge --NNLLNDKELENC FYLKANDLI I NRLKQN-----NSFIFSEVISHREPHDWSKTPVIYRASKQLFIK
#IMpn --NDTLNDPVL TGLFYLKANDVI IERLKQH-----HNFVFSSESFLHREPHDWSKTPVIYRASKQLFIK
#IMtu --DVTVADY-QGQHV F-DANAQIVRDLKTQSGPAAVNGPVLIRHETYEHPYPHCWRCRNPLIYRSVSSWFVR
#IBsu --TSEAPGF-EGMFYD-DANKAITQQLDEK-----GALVKLEFITHSYPHDWRTKKPTIFRATAQWFAS
#ICac --VDEVEPW-KGLFVK-KADPKILEYMKEN-----GTLYKSEKFTHSYPHCWRCDTPLLYYPRDSWFVR
#ISPn --MKNAGPEFEGQFY E-KVVPTVIEKLGNP-----PSCQEEISHSYPF DWRTKKPIIWRAPQWFAS
#ISyn --TEEAGKF-AGLNVLNDANEAI I NALQDQ-----KVL LKEEAYEHKYPYDWRTKKPTIFRATQWFI
#IPfl -GVYVPSLEFFGGQFIFKADQPI I EKLREV-----GALMQTAAIQH SYMHCWRHKTPLIYRATAQWFI
#ISmt EL PQSVRSIVRDEGLTKGRQVLD AETAKI I PCKLSDLNLLYKSHEYTHSYPYDWSKPKVI I RATPQWFAD
#ICmt -----LGHD-LDGKEV-LGEGQKIALRLLN-----HDIVHVS KHVHSYPYDWRTKKPVI IRSSEQWFID
#IHmt --TDVAGPE-LQNKAV-LEEGTDVVIKMLQ-----TAKNLLKEEKL VHSYPYDWRTKKPVVIRASKQWFI
#IHin --ISTTEFF-AGKGVF-EANPLVIEKLQEV-----GNLLKVEKIKHSYPHCWRHKTPI I FRATPQWFI
#IEco -LPGTYPTL-DGVNVF-KANDIVALLQEK-----GALLHVEKMQHSYPCCWRHKTPI I FRATPQWFVS
#INGo --ISETPRV-AGMSVW-EANPVILQWPEET-----GNLLASSKIEHSYAHWCWRHKTPLIYRATGQWFG
#ISce --TKDVPDF-EGVYVK-DADKLI I KYLTNT-----GNLLASQIRHSYPFCWRS DTPLLYRSVPWFVR
#ITth --TNVVSDF-AGVYIK-EADKLIRKNLKER-----GRL LVATTFTHSYPFCWRS DTPLIYKAVPSWFVR
#ICel --TSEVPDY-QGVYVK-DADKLI I KRLKEM-----GNLVRQAEVKHSYPFCWRS DTPLLYKAVPSWFIN
#INlo --NEEVPEY-KGMHV K-DADQSI I KDLGKK-----ILYEGKVYHRYPF CWRS DTPLIYKLVPNWVR
#IMth --TEDAGKY-RGLFVK-DADSDI I DDLRSK-----NLLLRAETI SHRYGFCWRCKTPIIYLATEQWFLK
#IMja --VE--GKW-KGVFVK-DADAEI I ETLKNK-----GLLVYAGKIKHSYPHCWRCKTPLLFRATEQWFLE
#IPfu --VE--GKW-KGVYVK-EADPQI I EHLKEK-----GYLVKAGEI EH KYPHCWRCKTPLIFRATDQWFLK
#IAfu --TEEAGKY-AGKHVK-EANDDI I DDLYRK-----DLLLAEERIVHRYGHCWRCKTPIIYRATEQWFIK
#ISac --TQDSGKY-AGKYVR-SASEEI I SDLKQR-----SALLHASKIVHRYPVCWRCKTPLILRAIEQWFIA

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#ITma  -----VDHNNLRQKVLEEIDKVKWIPEW-----GRNRIRSMVEERPDWCISRQRVWGTPIPAVKC
#IApy  MDIEYEGKTLSGES-LEEIEKVKWIPEY-----GKNRIKSMVENRPDWCISRQRFWGVPIITVFCY
#ICje  MDEPKLQGKTLREC-AKEQLLKTTTFYPQ----SGVKRIGSMVENRPDWCISRQRDWGTPIAFFRD
#IHpy  MDEPFIQNDGSQKT-LREVALDAIEKVEFVPSSGKNRLKTMIEENRPDWCLSRQRKWGVPLAFFID
#ISau  I-----SKVRQDI-LDAIENTNFKVNW-----GKTRIYNMVRDRGEWVISRQRVWGVPLPVFYA
#ISaR  -----TTNFKNEI-INNNNNIEWFPSH-----IKEGRMGNFLENMVDWNIGRNRYWGTPNLNVWIC
#ITpa  -----VEKIKDKM-LAANASICWQPSH-----IRDGRFGKWLVCARDWAISRDRYWGNPLPIWRC
#IBbu  -----VEKIKTKL-LEVNEKINWMPAH-----LKKGRFGKWLENAKDWAISRNRFWGNPIPIWIC
#ICtr  -----VEKVKSKM-LKANESIHWTPEH-----IKQGRFGKWLEGARDWAISRNRYWGTPIPIWRS
#IMge  -----TKSIKKQLKKQINQVNFLNSK-----NQLRLKEMLLQRDEWCISRQRVWGLPIPIVYA
#IMpn  -----TKSIQSK-LKRQIKRVKFVNN-----KNKERLQEMLLQRAEWCISRQRVWGLPIPLIYA
#IMtu  V-----TDFRDRM-VELNQQITWYPEH-----VKDGGQFGKWLQGARDWSISRNRYWGTPIPVWKS
#IBsu  I-----KDFRSDL-LDAIKETKWVPEW-----GEQRLHNMVRDRGDWCISRQRAWGVPIPVFYA
#ICac  M-----TSLRDKL-VENNNKIHWPDPN-----IRTGRFGKFVENVIDWGISRDRYWGTPLPIWQ
#ISpn  V-----SKFRQEILDEIEKVKFHSEW-----GKVRLYNMIRDGDWVISRQRAWGVPLPIFYA
#ISyn  V-----EGFRDQA-LKAIKEVTWIPTQ-----GENRITPMVGDRSDWCISRQRAWGVPIPVFYD
#IPfl  MDKEPTSGDTLRVRSLKAIEDTKFVPSW-----GQARLHSMIANRPDWCISRQRNWGVPIPFLLN
#ISmt  -----LHDVKNLALESISRVKFCPKR-----GYSRLSSFMSKSRNEWCISRQRSWGIPILSFYK
#ICmt  VD-----EIGKRASMLDDISVAAGDSD-----LRASLKQLVTTRKSWCISRQRVWGTPIPALVD
#IHmt  -----ITDIKTAAKELLKKVKFIPGS-----ALNGMVEMMDRRPYWCISRQRVWVPIPVFHH
#IHin  ME-----TQGLRQQALGEIKQVRWIPDW-----GQARIEKMVENRPDWCISRQRTWGVPMTLFVH
#IEco  MD-----QKGLRAQSLKEIKGVQWIPDW-----GQARIESMVANRPDWCISRQRTWGVPMSLFVH
#INGo  MDKAGSDGKTLRDKAIAVDDTEFFPPW-----GRARLESMEGRPDWVVSQRQRYWGTPMTFFVH
#ISce  V-----KNIVPQM-LDSVMKSHWVPNT-----IKEKRFANWIANARDWNVSRNRYWGTPIPLWVS
#ITth  V-----TALKDDL-LANNKKAYWVPKF-----AQEGRFNNWLQNVSDWCFSRSRFWGNPIPIWVS
#IHsa  V-----ENMVDQL-LRNNDLCYWVPEL-----VREKRFGNWLKDARDWTISRNRYWGTPIPLWVS
#ICel  V-----ETLIPRL-LANNDETYWVPAF-----VKEKRFANWLKDARDWAVSRNRFWGTPIPLWVS
#INlo  V-----KREIPRL-LESNSTINWIPES-----IGECHKFNWLSEARDWSISRNRFWGTPIPLWHC
#IMth  I-----TEIKDKM-LSELDRVQWIPSW-----AGESRFRNWIEANARDWTISRQRYWGIPPIWVC
#IMja  I-----SKIKDNI-IEHAKTVQWIPHW-----VETRYINGVKFVGDNWISRQRYWGIPVWVC
#IPfu  V-----SKVKEKIIKENDEKVTWYPEW-----VKIRFDNGVRDSDGWVISRQRYWGIPPLIWQS
#IAfu  I-----SELKDEM-LEEIDKVMWIPEW-----AGSARFKDWVSNAKDWCISRQRYWGIPVWIC
#ISac  V-----SKLKDHL-MGEIDRVRWIPDW-----GKTRIGNMVKEVRDWVISRQRFWGTPLPIWVC

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#ITma KECGEV-VLDPKVIIEHFMKIVEK-----EGTNAWFEKEVEELIPED
#IApy ENCGEV-IKDKEVFERIASLVEK-----HPGGTDVWFEEKSPEEILPEG
#ICje KNTKEV-IFDDELFDVFAAIFEK-----HGADAWWEFEIKDLIPTN
#IHpy KRTNKP-CFESEVLEHVANLF EK-----KGCDVWWEYSVKDLLPPS
#ISau EN-GEI-IMTKETVNHVADLFAE-----HGSNIWFEREAKDLLPEG
#ISaR ND-CNH-EYAPSSIKDLQNSIN-----KIDEDIELHRPYVDNI-
#ITpa VH-CGA-TDCIGSRTQLYERSGM-----LLEDLHKHVVDM-
#IBbu SK-TGK-KICIGSKKELENLSGQ-----KIEDLHKDQIDK-
#ICtr -D-DGE-LLVIGSIQELEALSGQ-----KIVDLHRHFIDE-
#IMge NN-KPL-LDFSTIQYTIKQLKKH-----GIDSWFEKDVTCTFL---
#IMpn DN-QPL-LDVTTIKYTIQQLKKY-----GIDSWFEKDINFFL---
#IMtu DD-PAY-PRIDVYGSLELERDF-----GVRPANLHRPYIDELTRP
#IBsu EN-GEP-VITDETIEHVSELFRQ-----HGSNIWFEKEAKDLLPEG
#ICac CE-CGH-RDCVGSIEELKEKGIN-----VPENIELHKPYIDEV
#ISPn ED-GTA-IMVAETIEHVAQLFEE-----HGSSIWWERDAKDLLPEG
#ISyn EETSEP-LLTEETINHVQQIIAE-----KGSDAWWELTVEELLPEA
#IPfl KESGELHPRTVELMEVVAQRVEQ-----QGIEAWFKLDAAELL---
#ISmt KSEPDSVLMNSEILAHAAIEKIKQ-----KGINAWFNDKDNMKEWL
#ICmt DNGGSYTSRKLIEWVAKLTRERE-----NTDVWWEIDVKEILENE
#IHmt KTKDEYLINSHTTEHIVKLVEQH-----GSDIWWTLPLLPKEVL
#IHin KETEELHPRTL DLLEEVAKRVER-----AGIQAWWDLDEKELL---
#IEco KDTEELHPRTLELMEEVAKRVEV-----DGIQAWWDLDAKEIL---
#INGo KETGELHPNSAELLEKVAQRIEE-----KGIEAWFSLDKSELL---
#ISce DDFEEV--VCVGSIEELEELT-----GVRNITDLHRDVIDKL-
#ITth EDFEEV--VCIGSVEELKKLT-----GATEITDLHKDFIDHL-
#IHsa DDFEEV--VCIGSVAELEELS-----GA-KISDLHRESVDHL-
#ICel EDGEEV--VCVGSIAELEELS-----GQ-KITDLHRESVDDV-
#INlo DG--KY--ICIGSIEELSRLS-----GR-KIDDIHRENVDDV-
#IMth EDCDSI--HVVGSI GELRELAV-----EGQLEGDFIHRPHVDRI-
#IMja EKCGKY--IVVGSVEELEEKMI-----NKDEVGEINDLHKPTVDKI-
#IPfu ED-GEI--YVVGSWKELVELTVAIEVN SERIDLPESYEEKLKVIEEKLGPEDLHRPYVDAF-
#IAfu EKCGEM--KVVGSINEIEWE-----NDLDLHRPKIDAV-
#ISac SNCQNI--IVVGGVDELGKIPI-----NQVPQDLHRPWIDSV-

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#ITma FVCPKCGKRSFEKMLDITLDVWIDSGASFEYIT-----TKREDHPFPLDMYLEGSDQHRG
#IApy YKCPKCGGTSFRKEEDILDVWFDSGCSHASV-----IRPLGFEEKADLYLEGSDQHRG
#ICje ---SKYNAENLEKVYDILDVWFDSGSTFNAVL-----NSGLYDAGEKRASMYLEGSDQHRG
#IHpy ---YQEDAKHYEKIMHILDVWFDSGSTFKAVL-----EDYHGEKGQSPSDVILEGSDQHRG
#ISau FTHPGSPNGTFTKETDIMDVWFDSGSSHRGVL-----ETRPELSFPADMYLEGSDQYRG
#ISaR TLSCPCKCNGKMSRVEEVIDVWFDSGSMFPAQHHPFD-NQKIFNQHFPADFIAEGVDQTRG
#ITPa VTIPCACGSVMRRVPEVLDCWFESGAMPYAQQHYHYPFE-HATDFERYFPAHFISEGLDQTRG
#IBbu ITWPSKDGKGFIRTSEVLDCWFESGAMPYASNHYHYPFE-NEINFKNIFPADFIAEGLDQTRG
#ICtr -IEINQNGKSFRRIPIYVFDVWFDSGAMPYAQNHYHYPFE-RAEETEACFPADFIAEGLDQTRG
#IMge KPDKTKKWKVYHKEIDITLDVWFDSGSSYNVL-----EINKYGSIAADLYIEGSDQYRG
#IMpn NPKKIQPGVEYKKETDTLEVWFDSGSTYNVL-----ISNKLNFADLYLEGSDQYRG
#IMtu NPDDPTGRSTMRRIIPDVLVWFDSGSMFYAQVHYHYPFE-NLDWFQGHYPGDFIVEYIGQTRG
#IBsu FTHPGSPNGTFTKEQDIMDVWFDSGSSHQAVL-----EERDDLVRPADLYLEGSDQYRG
#ICac KLTCPKCGKPMTRTEEVIDVWFDSGSMFPAQLHYHYPFE-NKEVFENTFPAQFISEAVDQTRG
#ISpn FTHPGSPNGEFKKETDIMDVWFDSGSSWNGVV-----VNRPELTYPADLYLEGSDQYRG
#ISyn ---YRNNGRTYRKGEDTMDVWFDSGSSWAAVA-----NAKNRPLKYPVDMYLEGSDQHRG
#IPfl ----GDEAPLYDKISDTLDVWFDSGTTHWHVL---RGSHPMGHETGPRADLYLEGSDQHRG
#ISmt PEKYHDVAHEYCRSQDTMDVWFDSGSSWSVIKDFYEKSLKLSKLPSPLYQVCLEGSDQHRG
#ICmt -----EKNTDIMDVWLDGLAWHAAR-----DNDTEREHVADVLEGVDFQFRG
#IHmt SEVGGPDALYVPGQDILDIWFDSGTWSYV-----LPGPDQRADLYLEGKDQLGG
#IHin ----GADAETYRKVPDTLDVWFDSGSTYSSVV-----ANRLEFNQDIDMYLEGSDQHRG
#IEco ----GDEADQYVKVPDTLDVWFDSGSTHSSVV-----DVRPEFAGHAADMYLEGSDQHRG
#INGo ---SAEDCEHYDKLPDTMDVWFDSGSTHYSVV-----KQREELEWPADLYLEGSDQHRG
#ISce TIPSKQKGKDLKRIEEVFDCWFESGSMFYASQHYHYPFE-NTEKFDERVPANFISEGLDQTRG
#ITth TIPSKQKGKGLRRIDEVFDCWFESGSMFYAQVHYHYPFE-NKREFEDAFPADFIAEGIDQTRG
#IHsa TIPSRGCGKSLHRISEVFDCWFESGSMFYAQVHYHYPFE-NKREFEDAFPADFIAEGIDQTRG
#ICel TIPSRSGRGVLKRVSEVFDCWFESGSMFYAQNHYPFE-NRKIFEDNFPADFIAEGIDQTRG
#INlo VICKDGEK--YRIIEVFDCWFESGCMFYAQRHWPFE----CDNLCLPADFVAEGVDQTRG
#IMth ILECGRCGGRMKRTPDVLVWIDSGVAGWAALHYPRE--KELFSEWFPYDFITEGHDQTRG
#IMja KLRCE--CGGEMKRVDPDVLVWFDSGLAPYASIGV-----KELKKADFITEGHDQVTK
#IPfu IIKVN--GKEMRRVKDVVDVWFDSGIASWASLGYPR--NKELFEKLWPADFIVEGEDQVTK
#IAfu TFSCQ--CGGVMRRVPDVFVWFDSGVASWGSIAAYPL--RKDKFEELWPADFITEGHDQTRG
#ISac VLDVKNVEGQARRISDVADVWFDSGVAFASLGQDW--RKR--WSELGPVDLVLEGHDLRG

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#ITma WFHSSIFLAVAKRGSAPYKEVLT--HGFIKDEQGR-----
#IApy WFQASLLESVGSYGEAPYRSVLT--HGFIVDEQGR-----
#ICje WFQSSLLVGTAINESTPYESILT--HGFTTDEKGQ-----
#IHpy WFQSSLLIGCVLNNQAPFKKVIT--HGFIVDEKGE-----
#ISau WFNSSITTSVATRGVSPYKFLLS--HGFVMDGEGK-----
#ISaR WFYSLLVISTILKGKSSYKRALS--LGHILDSNGK-----
#ITpa WFYTLTILAVALFERPAFENCIV--TGLVLASDGK-----
#IBbu WFYTLTILGTALFENTAFKNVIV--NGLVLSSDGR-----
#ICtr WFYTLTVIAAALFDQPAFKNVIV--NGIILAEDGN-----
#IMge WFNSSSNCGIIQNDLIPFKSLVS--HGFTLDENG-----
#IMpn WFNSSASCGIIQTDQLPFKSLIS--HGFTLDEHGN-----
#IMtu WFYTLHLVLATALFDRPAFKTCVA--HGIVLGFQDGQ-----
#IBsu WFNSSLSTAVAVTGKAPYKGVLS--HGFALDGEGR-----
#ICac WFYTLAISTSIQDKSSFENCIV--LGHVLDKHGL-----
#ISpn WFNSSLITSVANHGVAPYKQILS--QGFDLGEKGE-----
#ISyn WFQSSLLTSVAVNGIAPYKTVLT--HGFVLDEKGH-----
#IPfl WFHSSLLTGCAIDNHAPYRELLT--HGFTVDETR-----
#ISmt WFQSSLLTKVASSNVPVAPYEEVITHGFTLDENGL-----
#ICmt WFQSSLLTSVAVQNKLPHYKIIIV--HGFCIDENNN-----
#IHmt WFQSSLLTSVAARKRAPYKTVIV--HGFTLGEKGE-----
#IHin WFMSSMLSTATDSKAPYKQVLT--HGFTVDGQGR-----
#IEco WFMSSLMISTAMKGKAPYCQVLT--HGFTVDGQGR-----
#INGo WFQSSMLTGCASSMGRAPYKQLLT--HGFVVDQNGR-----
#ISce WFYTLAVLGTHLFGSVPYKNVIV--SGIVLAADGR-----
#ITth WFYTLNVISTALRNSNPYKNLIV--NGIVLAADGK-----
#IHsa WFYTLVLATALFGQPPFKNVIV--NGLVLASDGQ-----
#ICel WFYTLVLSTALFNKPPFKNLIC--NGLVLASDGA-----
#INlo WFYTMHVISTVLLAKPRFRNCIV--NGIVLASDKK-----
#IMth WFYSQLGCGVIALDETPYRRVLM--HGFTLDEEGR-----
#IMja WFYSQHALSAIVFNDIPYKKCLM--HGFTLDEHGD-----
#IPfu WFYSQQAASVIAFDTVPYRAVAM--HGYVLDEKGD-----
#IAfu WFYSQLGTSVVCFDKAPYKAVLM--HGFTLDEQGR-----
#ISac WFFSLLRTGVILMDKAPYEAVLV--HGFMLDEQGREMHKSSGNYVEPSQVVSKYGQRYVKIMV

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#ITma -----KMSKSL
#IApy -----KMSKSL
#ICje -----KMSKSK
#IHpy -----KMSKSK
#ISau -----KMSKSL
#ISaR -----KMSKSK
#ITpa -----KMSKAL
#IBbu -----KMSKSF
#ICtr -----KMSKRL
#IMge -----KMSKSL
#IMpn -----KMSKSL
#IMtu -----KMSKSL
#IBsu -----KMSKSI
#ICac -----KMSKHK
#ISpn -----KMSKSL
#ISyn -----KMSKSL
#IPfl -----KMSKSL
#ISmt -----KMSKLV
#ICmt -----KMSKSI
#IHmt -----KMSKSL
#IHin -----KMSKSI
#IEco -----KMSKSI
#INGo -----KMSKSI
#ISce -----KMSKSL
#ITth -----KMSKSK
#IHsa -----KMSKRK
#ICel -----KMSKSK
#INlo -----KMSKRL
#IMth -----KMSKSL
#IMja -----KMSKSL
#IPfu -----KMSKSL
#IAfu -----KMSKSL
#ISac TLEIPHGKMSKSH